Amendments to the Specification

Please replace the present title with the following amended title:

-- PROCESS FOR THE PURIFICATION OF RECOMBINANT ALBUMIN --

Please replace the three one sentence paragraphs beginning on page 24, line 22, and ending on page 25, line 2, with the following amended paragraphs:

~~Figures~~ Figure 10 (SEQ ID NO. 1) and 11 (SEQ ID NO. 2) represent two DNA sequences with homology to the protein encoding region *Saccharomyces cerevisiae PMT1*.

~~Figures~~ Figure 12 (SEQ ID NO. 3), Figure 13 (SEQ ID NO. 4), Figure 14 (SEQ ID NO. 5), and Figure 15 (SEQ ID NO. 6) ~~to 15~~ represent four DNA sequences with homology to the protein encoding region *Saccharomyces cerevisiae PMT7.*

~~Figures~~ Figure 16 (SEQ ID NO. 7) and Figure 17 (SEQ ID NO. 8) represent two DNA sequences with homology to the protein encoding region *Saccharomyces cerevisiae PMT5*.

{Remainder of page blank}

Please replace the paragraph beginning on page 32, line 18 (under "Example 1"), and ending on page 33, line 3 with the following amended paragraph:

The cloning strategy for construction of the albumin-producing micro-organism was as disclosed in EP 431 880 except that the 3′ end of the albumin coding sequences and its junction with the *ADH1* transcription termination sequence were altered such that the ADH coding sequence was eliminated and such that two consecutive in-frame translation stop codons were present, followed by a third stop codon downstream, as follows:

.....	L	G	L	stop	stop	A	stop	
.....	TTA	GGC	TTA	TAA	TAA	GCT	TAA	.....

SEQ ID NO. 9

Please replace the paragraph on page 33, lines 4 to 21 with the following amended paragraph:

This was achieved by modification of the *ADH1* terminator from plasmid pAYE309, described in EP 431 880, by PCR mutagenesis using two single stranded oligonucleotides, JMADH1 and JMADH2 with the sequences:

JMADH1

SEQ ID NO. 10

*Hin*dIII

5′ – GCATAAGCTTTGGACTTCTTCGCCAGAGGTTTGGTCAAG – 3′

JMADH2

SEQ ID NO 11

*Not*I *Bam*HI
3'-T GGACAACATTAGCAAGAAGGTGTGCCTAGCGCCGGCGCCTAGGTACG-5'

Please replace the paragraph beginning on page 33, line 24 and ending on page 34, line10, with the following amended paragraph:

The PCR conditions were 25 cycles of 94°C for 60 seconds, 37°C for 120 seconds and 72°C for 180 seconds. The 0.48kb PCR product was digested with both *Hin*dIII and *Bam*HI and ligated into plasmid pBST+, described in WO 97/24445, similarly digested with *Hin*dIII and *Bam*HI, to create plasmid pAYE440 (Fig. 2). The *ADH1* terminator was further modified by PCR mutagenesis using two single stranded oligonucleotides, AT19R and the universal - 40 primer with the sequences:

AT19R

SEQ ID NO. 12

*Hin*dIII
5' - AGTCCAAGCTTAATTCTTATGATTTATGAT - 3'

-40

SEQ ID NO. 13

3' - CAGCACTGACCCTTTTG - 5'.

Please replace the paragraph beginning on page 35, lines 1-18, with the following amended paragraph:

The double stranded oligonucleotide linker, AT21/AT22 was ligated into *Afl*III/*Hind*III cut pDB2241 and comprised an *Afl*III site at its 5N end, a stuffer region and then the *Bsu*36I to *Hind*III sequence of the HSA coding DNA, but with the addition of an extra TAA translation stop codon. Clones with the linker inserted were checked by DNA sequencing and the correct plasmid designated pDB2242 (Fig. 5).

Linker AT21/22

AT21

*Afl*III *Bsu*36I *Hind*III

SEQ ID NO. 14

TTA AGA GTC CAA GCC TTA GGC TTA TAA TA

CT CAG GTT CGG AAT CCG AAT ATT ATTCGA

SEQ ID NO. 15

A L G L Stop Stop

SEQ ID NO. 16

Please insert Pages 1 to 5 of the enclosed Sequence Listing after page 84 of the application as filed (after the specification and before Figure 1 pursuant to 37 CFR. §1.77(b)).